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AMENDMENTS TO THE CLAIMS

Claims 1 to 20 (Cancelled)

- 21. (New) An isolated polypeptide comprising a polypeptide sequence selected from the group consisting of:
 - (a) an isolated polypeptide comprising amino acids 1 to 541 of SEQ ID NO:2; and
 - (b) an isolated polypeptide comprising amino acids 2 to 541 of SEQ ID NO:2.
 - 22. (New) The isolated polypeptide of Claim 21, wherein said polypeptide is (a).
 - 23. (New) The isolated polypeptide of Claim 21, wherein said polypeptide is (b).
 - 24. (New) An isolated polypeptide produced by a method comprising:
- (a) culturing an isolated recombinant host cell comprising a vector that comprises the coding region encoding the polypeptide of Claim 21 under conditions such that the polypeptide of Claim 21 is expressed; and
 - (b) recovering said polypeptide.
- 25. (New) The isolated polynucleotide of Claim 21 wherein said polypeptide sequence further comprises a heterologous polypeptide sequence.
- 26. (New) The isolated polynucleotide of Claim 25 wherein said heterologous polypeptide is the Fc domain of an immunoglobulin.
- 27. (New) An isolated polypeptide comprising the polypeptide encoded by the BGS-42 cDNA clone A in ATCC Deposit No. PTA-4454.
- 28. (New) An isolated polypeptide comprising the polypeptide encoded by the BGS-42 cDNA clone B in ATCC Deposit No. PTA-4454.
- 29. (New) An isolated polypeptide comprising the polypeptide encoded by the BGS-42 cDNA clone C in ATCC Deposit No. PTA-4454.
- 30. (New) An isolated polypeptide comprising a polypeptide sequence that is at least 95.0% identical to amino acids 2 to 541 of SEQ ID NO:2, wherein percent identity is calculated using a CLUSTALW global sequence alignment using default parameters, and wherein said polypeptide has tubulin tyrosine ligase activity.
- 31. (New) An isolated polypeptide consisting of at least 50 contiguous amino acids of SEQ ID NO:2.
 - 32. (New) An isolated polypeptide comprising amino acids 73 to 365 of SEQ ID NO:2.

- 33. (New) An isolated polypeptide comprising amino acids 133 to 374 of SEQ ID NO:2.
- 34. (New) An isolated polypeptide comprising amino acids 2 to 541 of SEQ ID NO:2, wherein the amino acid located at amino acid position 515 is a glutamic acid.
- 35. (New) An isolated polypeptide comprising amino acids 2 to 541 of SEQ ID NO:2, wherein the amino acid located at amino acid position 524 is a serine.
- 36. (New) An isolated polypeptide comprising at least 394 contiguous amino acids of SEQ ID NO:2.
- 37. (New) The isolated polypeptide of Claim 21(a) or (b), wherein said polypeptide has tubulin tyrosine ligase activity.
- 38. (New) The isolated polypeptide of Claim 21(a) or (b), wherein said encoded polypeptide has one amino acid substitution.
- 39. (New) An isolated polypeptide comprising a polypeptide encoded by a polynucleotide that hybridizes under stringent conditions to the polynucleotide encoding amino acids 2 to 541 of SEQ ID NO:2, wherein said stringent conditions are as follows: an overnight incubation at 42 degree C in a solution comprising 50% formamide, 5x SSC (750 mM NaCl, 75 mM trisodium citrate), 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and 20 μg/ml denatured, sheared salmon sperm DNA, followed by washing the filters in 0.1x SSC at about 65 degree C, wherein said polynucleotide encodes a polypeptide that has tubulin tyrosine ligase activity.
 - 40. (New) An isolated polypeptide produced by a method comprising:
- (a) culturing an isolated recombinant host cell comprising a vector that comprises
 a coding region operatively linked to nucleotides -2057 to -1 of the sequence provided in Figures
 7A-B under conditions such that a polypeptide is expressed; and
 - (b) recovering said polypeptide.